A SWLA1: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VL DOMAIN OF CHIMERIC ANTIBODY TEDW

EcoRV (242) GGGGATATCCACCATGGAGACAGACACTCCTGCTATGGGTGCTGCTGCTCTGGGTTCCAGGTTCCACAGGTGACATTGT ▶ METDTLLLWVLLLWVPGSTGDIV Pstl (377) GCTGACCCAATCTCCAGTTTCTTTGGCTGTGTCTCTAGGGCAGAGGCCACCATATCCTGCAGAGCCAGTGAAAGTGTTGA LTQSPVSLAVSLGQRATISCRASESVD Kpnl (427) TAGTTATGGCAATAGTTTTATGAACTGGTACCAGCAGAAACCAGGACAGCCACCCCAACTCCTCATCTATCGTGCATCCAA SYGNSFMNWYQQKPGQPPQLLIYRASN Xbal (482) TCTAGAATACGGGATCCCTGCCAGGTTCAGTGGCAGTGGGTCTAGGACAGACTTCACCCTCACCATTAATCCTGTGGAGGC LEYGIPARFSGSGSRTDFTLTINPVEA TGATGATGTTGCAACCTATTACTGTCAGCAAAATAATGCGGATCCTCCCACGTTCGGAGGGGGGACCAAGTTGGAAATCAA D D V A T Y Y C Q Q N N A D P P T F G G G T K L E I K Sall (650) ACGTAAGTCGACGCT RKS

B SWLA1: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VH DOMAIN OF CHIMERIC ANTIBODY TEDW

EcoRV (242)

SWLA2: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VL DOMAIN OF CHIMERIC ANTIBODY TEFE

E

RV (243)

GGGGATATCCACCATGGATTTTCAAGTGCAGATTTTCAGCTTCCTGCTAATCAGTGTCACAGTCATATTGACCAATGGAGAAA

▶ M D F Q V Q I F S F L L I S V T V I L T N G E

Pstl (384) BstEll (372)

TTTTGCTCACCCCGTCTCCAGCAATCATAGCTGCATCTCCTGGGGAAAAGGTCACCATCACCTGCAGTGCCAGCTCAAGTGTT

FILL TPSPAIIAASPGEKVTITCSASSSV

Kpnl (419)

AGTTACATGAACTGGTACCAGCAGAAACCAGGATCTTCCCCCCAAAATCTGGATTTATGGTGTATCCAACCTGGCTTCTGGAGT F S Y M N W Y Q Q K P G S S P K I W I Y G V S N L A S G V TCCTGCTCGCTTCAGTGGCAGTGGGTCTGGGACATCTTTCTCTTTCACAATCAACAGCATGGAGGCTGAAGATGTTGCCACTT

▶ PARFSGSGSGTSFSFTINSMEAEDVAT

ATTACTGTCAGCAAAGGAGTAGTTACCCATTCACGTTCGGCTCGGGGACCAAGCTGGAAATAAAACGTAAGTCGACGCT PY Y C Q Q R S S Y P F T F G S G T K L E I K R K S

SWLA2: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VH DOMAIN OF CHIMERIC ANTIBODY TEFE

EcoRV (242)

Ndel (295)

GGGATATCCACCATGGACAGGCTTACTTCTTCATTCCTGCTACTGATTGTTCCTGCATATGTCCTCTCCCAGGTTACTCT

▶ M D R L T S S F L L L I V P A Y V L S Q V T L

KESGPGILQPSQTLSLTCSFSGFSLR

▶TYGIGVGWIRQPSGRGLEWLAHIWWND

Scal (484) AATAAGTACTATAACACAGTCCTGAAGAGCCGGCTCACAATCTCCAAGGATACCTCCAACAACCAGGTATTCCTCAAGAT ▶ N K Y Y N T V L K S R L T I S K D T S N N Q V F L K I

CGCCAGTGTGGACACTGCAGATACTGCCACATACTACTGTGCGCGAATAGAGGGGGGGCTCGGGCTACGATGTTATGGACT

A S V D T A D T A T Y Y C A R I E G G S G Y D V M D

Nhel (675)

Sall (696)

ACTGGGGTCAAGGAATCTCAGTCACCGTCTCTTCAGCTAGCACAACACCCCCATCTGTCGACCCA YWGQGISVTVSSAS

A SWLA3: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VL DOMAIN OF CHIMERIC ANTIBODY TEFC

EcoRV (242)

GGGATATCCACCATGATGAGTCCTGCCCAGTTCCTGTTTCTGTTAGTGCTCTGGATTCGGGAAACCAACGGTGATGTTGTG

MMSPAQFLFLLVLWIRETNGDVV

BstEl(347)

ATGACCCAGACTCCACTTTGTCGGTTACCATTGGACAACCAGCCTCCATCTCTTGCAAGTCAAGTCAGAGCCTCTTA

DRDGRTYLSWLLQRPGQSPKRLIYLVS

AAACTGGACTCTGGAGTCCCTGACAGGTTCACTGGCAGTGGATCAGGGACAGATTTCACACTGAAAATCAGCAGAGTGGAG

K L D S G V P D R F T G S G S G T D F T L K I S R V E GCTGAGGATTTGGGGAGTTATTATTGCTGGCAAGGTACACATTTTCCGCTCACGTTCGGTGCTGGGACCAAGCTGGAGCTG

► A E D L G V Y Y C W Q G T H F P L T F G A G T K L E L Sall (653)

AAACGTAAGTCGACC

KRKS

B SWLA3: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VH DOMAIN OF CHIMERIC ANTIBODY TEFC

EcoRV (1425)

GATATCCACCATGGACTTCGGGTTGAGCTTGGTTTTCCTTGTCCTTACTTTAAAAGGTGTCCAGTGTGACGTGAAGCTGGT

GGGTCAAGGAACCTCAGTCACCGTCTCTTCAGCTAGCTCAACACCCCCATCAGTCGACCCA

SWLA1: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE ABERRANT VL DOMAIN

EcoRV ▶ M E T D T L L L W V L L L W V P G TCCACTGGTGACATTGTGCTGACACAGTCTCCTGCTTCCTTAGCTGTATCTCTGGGGCAGAGGGCCACCATCTCATAC STGDIVLTQSPASLAVSLGQRATISY AGGGCCAGCAAAAGTGTCAGTACATCTGGCTATAGTTATATGCACTGGAACCAACAGAAACCAGGACAGCCACCCAGA PRASKSVSTSGYSYMHWNQQKPGQPPR EcoO1091 CTCCTCATCTATCTTGTATCCAACCTAGAATCTGGGGTCCCTGCCAGGTTCAGTGGCAGTGGGTCTGGGACAGACTTC LLIYLVSNLESGVPARFSGSGSGTDF PfIMI ACCCTCAACATCCATCCTGTGGAGGAGGAGGATGCTGCAACCTATTACTGTCAGCACATTAGGGAGCTTACACGTTCG T L N I H P V E E E D A A T Y Y C Q H I R E L T R S GAGGGGGGACCAAGCTGGAAATAAAACGGNCTNATGCTGCACCAACTGTATCCATCTTNAAAANCATCAGTTCTAGAG ▶ E G G P S W K • **EcoRI** AAGGGCGAATTCC

FIG. 5

SWLA1: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE NON-EFFECTIVE 2ND VH DOMAIN

EcoRV (242)

CIGGICCGCTIACIGGGGCCAAGGGACIGIGGICACIGICICTICAGCIAGCACACACCCCCATCAGICIACCCA

W S A Y W G Q G T V V T V S S A S

SWLA1: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE ABERRANT VH DOMAIN

EcoRV EcoRI ▶ M E T D T L L L W V L L L W V P G ▶ STGDIVLTQSPASLAVSLGQRATISY ▶ R A S K S V S T S G Y S Y M H W N Q Q K P G Q P P R EcoO1091 LLIYLVSNLESGVPARFSGSGSGTDF ACCCTCAACATCCATCCTGTGGAGGAGGAGGATGCTGCAACCTATTACTGTCAGCACATTAGGGAGCTTACACGTTCG ▶T L N I H P V E E E D A A T Y Y C Q H I R E L T R S GAGGGGGGACCAAGCTGGAAATAAAACGGNCTNATGCTGCACCAACTGTATCCATCTTNAAAANCATCAGTTCTAGAG FEGGPSWK • **EcoRI** AAGGGCGAATTCC

FIG. 7

SWLA2: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE ABERRANT VH DOMAIN

EcoRI EcoRV

GGAATTCGCCCTTGGGGATATCCACCATGGGATGGAGCTGGGTCATGCTCTTTCTCCTGGCAGGAACTGCAGGTGTCCT

M G W S W V M L F L L A G T A G V L

EcoRV

CTCTGAGGTCCAGCTGCAACAGTCTGGACCTGAGCTGGTGAAGCCTGGGGCTTCAGTGAAGATATCCTGCAAGACTTCT

SEVQLQQSGPELVKPGAACACATGCACTGGGTGAAACAGAGCCATGGAAAGAGCCTTGAGTGGATTGAGGGTATTA

GGATACACATTCACTGAATACAACATGCACTGGGTGAAACAGAGCCATGGAAAGAGCCTTGAGTGGATTGAGGTATTA

GYTFTEYNMHWVVKQSHGKSLEWIGGI

ATCCTAACAATGGTGGTACTAGTTACAACCAGAAGTTCAAGGCCAAGGCCACATTGACTGTAGACAAGTCCTCCAGCAC

NPNNGGTSYNQKFKAAKATTCTGCAGCTCTTTACTGTGCAAGGGGGGTTTATGATGGTTA

AGCCTACATGGAGCTCCGCAACCTGACATCTGAGGATTCTGCAGTCTATTACTGTGCAAGGGGGGTTTATGATGGTTA

AYMELRNLTSEDSAVYYCARGCCCAACAGCCCCATCGGTCTATTCCAC

SLLTTGAAKACAACAGCCCCATCGGTCTCTCACAGTCTCCTCAGCCAAAACAACAGCCCCATCGGTCTATTCCAC

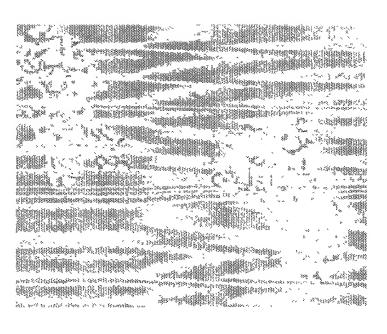
SLLTTGAAKACAACAGCCCCATCGGTCTATTCCAC

WPL

TGGCCCCTG

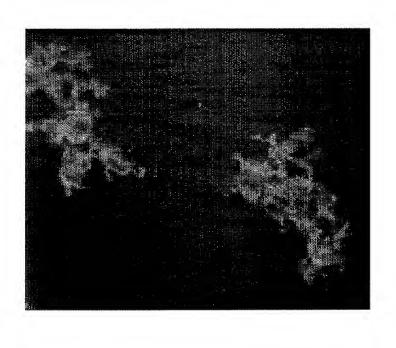
WPL

LIGHT AND FLUORESCENT MICROSCOPE IMAGES CHIMERIC ANTIBODY TEDW BINDING TO S. MUTANS



S. mutans +TEDW

Light microscope



+Sigma F9512 S. mutans +TEDW



+Sigma F5387 +TEDW

Fluorescent microscope